

Analysis of the Constituents in Rat Serum after Oral Administration of Fufang Zhenzhu Tiaozhi Capsule by UPLC–Q–TOF–MS/MS

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Abstract A rapid and sensitive UPLC/Q–TOF–MS method has been established for analysis of the constituents in rat serum after oral administration of Fufang Zhenzhu Tiaozhi (FTZ) capsule, an effective compound prescription for treating hyperlipidemia in the clinic. The UPLC/MS information of samples was obtained first in FTZ preparation and FTZ-treated rat serum. Mass spectra were acquired in both negative and positive ion modes. Thirty-six constituents in rat serum after oral administration of FTZ were detected, including the alkaloids, ginsenosides, pentacyclic triterpenes, and their metabolites. These chemicals were identified based on the retention time and mass spectrometry data with those of authentic standards or comparison of the literatures reports. Twenty-seven prototype components originated from FTZ and nine were the metabolites of the FTZ constituents. These results shed light on the potential active constituents of the complex traditional Chinese medicinal formulas.

Keywords UPLC/Q–TOF–MS · Serum pharmacology · Metabolite · Fufang Zhenzhu Tiaozhi capsule

Introduction

Fufang Zhenzhu Tiaozhi (FTZ) capsule, the patentable Chinese herbal medicine prescription, including *Rhizoma Coptidis*, *Radix Salvia Miltiorrhiza*, *Radix Notoginseng*, *Fructus Ligustri Lucidi*, *Herba Cirsii Japonici*, *Cortex Eucommiae*, *Fructus Citri Sarcodactylis* and *Radix Atractylodes Macrocephala*. FTZ has been prescribed for 12 years by virtue of the potential to regulate abnormal lipid metabolism for treatment of dyslipidemia, atherosclerosis, and related disease [1, 2]. Clinical practice on more than 3,000 dyslipidemic patients demonstrated that FTZ is very safe and less harmful side effects [3, 4]. Giving FTZ not only markedly decrease the levels serum total cholesterol, glycerinate and low-density lipoprotein cholesterol while raising high-density lipoprotein cholesterol, but also improves hepatic tissue pathologic states, and prevents atherosclerosis [5, 6].

At present, hundreds of constituents have been identified, respectively and systematically, from the herbs that compose FTZ [7]. Constituents such as oleanolic acid, salvianolic acid A, salvianolic acid B, notoginsenoside R₁, ginsenoside Rb₁, ginsenoside Rg₁, berberine, palmatine and jateorhizine have been experimentally verified [8, 9]. However, it remains unclear as to which constituents are responsible for the lipid-modulating functions of the drug; furthermore, there has been no integrated study of the constituents of the formula which is not simply a blend of the individual herbs but an integrated prescription.

Serum pharmacochimistry, which is an experimental technique focusing on the analysis of serum samples obtained after dosing, is based on the hypothesis that most effective constituents need to be absorbed into the blood to elicit activities after administration of traditional Chinese medicines (TCMs), and the components absorbed and

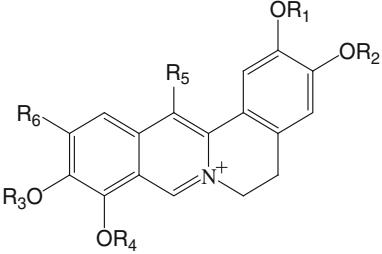
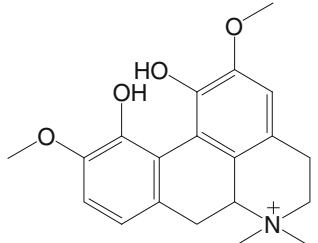
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Table 1 Structures of the Rhizoma Coptidis constituents identified in FTZ preparation and serum samples from FTZ-treated rats

Type	Substituent group	Name of constituent	MW	Observed
	$R_1=R_2=R_3=R_4=CH_3, R_5=H, R_6=OH$	Hydroxyl-palmatine	368	In serum and in preparation
	$R_1+R_2=CH_2, R_3=H, R_4=CH_3, R_5=R_6=H$	Thalifendine	322	In serum and in preparation
	$R_1=H, R_2=R_3=R_4=CH_3, R_5=R_6=H$	Columbamine	338	In serum and in preparation
	$R_1=R_2=CH_3, R_3+R_4=CH_2, R_5=R_6=H$	Epiberberine	336	In serum and in preparation
	$R_1+R_2=R_3+R_4=CH_2, R_5=R_6=H$	Coptisine	320	In serum and in preparation
	$R_1=CH_3, R_2=H, R_3=R_4=CH_3, R_5=R_6=H$	Jatrorrhizine	338	In serum and in preparation
	$R_1+R_2=CH_2, R_3=CH_3, R_4=H, R_5=R_6=H$	Berberrubine	322	In serum and in preparation
	$R_1=R_2=R_3=R_4=CH_3, R_5=R_6=H$	Palmatine	352	In serum and in preparation
	$R_1+R_2=CH_2, R_3=R_4=CH_3, R_5=R_6=H$	Berberine	336	In serum and in preparation
	$R_1=R_2=R_3=R_4=CH_3, R_5=CH_3, R_6=H$	Dehydrocorydaline	366	In serum and in preparation
	$R_1+R_2=CH_2, R_3=R_4=CH_3, R_5=CH_3, R_6=H$	13-Methylberberine	350	In serum and in preparation
	$R_1=R_2=R_3=R_4=CH_3, R_5=CH_2CH_3, R_6=H$	13-Ethyl-5,6-dihydro-2,3,9,10-tetramethoxy-dibenzo[a,g]quinolizinium	380	In preparation only
	$R_1=CH_3, R_2=H, R_3=R_4=CH_3, R_5=CH_3, R_6=H$	Dehydrocorybulbine	352	In preparation only
	$R_1=CH_3, R_2=GlcUA, R_3=R_4=CH_3, R_5=R_6=H$	Jatrorrhizine 3-O- β -D-glucuronide	514	In preparation only
			Magnoflorine	342

In serum and in preparation: the constituent was observed both in FTZ-treated serum and FTZ preparation, in preparation only: the constituent was only observed in FTZ preparation, in serum only: the constituent was only observed in FTZ-treated serum

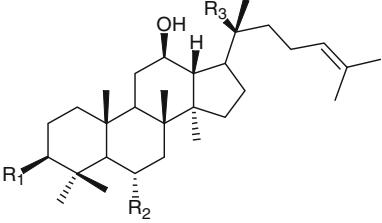
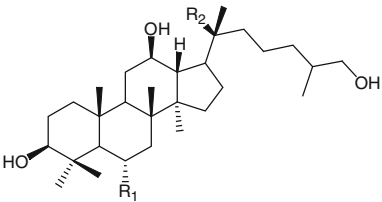
MW molecular weight

metabolites formed can be determined simultaneously in order to identify the in vivo active forms from TCM formulas [10–14]. On the other hand, the rapid development of analytical techniques, such as UPLC coupled with HDMS technique in recent years provide a powerful tool for qualitative and quantitative analysis of complicated samples such as TCMs [15, 16].

The present study examined the constituents of rat serum after oral administration of FTZ using combined

UPLC/Q–TOF–MS/MS. From a comprehensive analysis of a FTZ preparation, rat serum collected from FTZ-treated group and control group, 27 prototype components, and nine metabolites originating from FTZ were identified. To the best of our knowledge, this is the first systematical study on identifying the possible effective constituents in FTZ. The information will guide us to explore the mechanism under the lipid-modulating effect of FTZ in the following investigation.

Table 2 Structures of the Radix Notoginseng constituents identified in FTZ preparation and serum samples from FTZ-treated rats

Type	Substituent group	Name of constituent	MW	Observed
	$R_1=OH, R_2=OGlc(2-1)Xyl, R_3=OGlc$	Notoginsenoside R_1	932	In serum and in preparation
	$R_1=OH, R_2=OGlc, R_3=OGlc$	Ginsenoside R_{g1}	800	In serum and in preparation
	$R_1=OH, R_2=OGlc(2-1)Rha, R_3=OGlc$	Ginsenoside R_e	946	In preparation only
	$R_1=OH, R_2=OGlc, R_3=OH$	Ginsenoside R_{h1}	638	In serum and in preparation
	$R_1=OH, R_2=OH, R_3=OGlc$	Ginsenoside F_1	638	In serum and in preparation
	$R_1=OGlc(2-1)Glc, R_2=H, R_3=OGlc(6-1)Glc$	Ginsenoside R_{b1}	1108	In serum and in preparation
	$R_1=OGlc(2-1)Glc, R_2=H, R_3=OGlc$	Ginsenoside R_d	946	In serum and in preparation
	$R_1=OH, R_2=OH, R_3=OH$	Protopanaxatriol	476	In serum only
	$R_1=OGlc(2-1)Glc, R_2=H, R_3=OH$	Ginsenoside R_{g3}	784	In serum only
	$R_1=OGlc, R_2=OH$	25-Hydroxy-ginsenoside R_{h1}	656	In serum only
	$R_1=OH, R_2=OGlc$	25-Hydroxy-ginsenoside F_1	656	In serum only

In serum and in preparation: the constituent was observed both in FTZ-treated serum and FTZ preparation, in preparation only: the constituent was only observed in FTZ preparation, in serum only: the constituent was only observed in FTZ-treated serum

MW molecular weight

Experimental

Chemicals and Materials

Authentic standards such as chloramphenicol, danshensu, protocatechuic acid, protocatechuic aldehyde, salidroside, rosmarinic acid, salvianolic acid B, specnuezhenide, salvianolic acid A, jatrorrhizine, notoginsenoside R_1 , palmatine, berberine, ginsenoside R_{g1} , ginsenoside R_e , 5,7-dimethoxycoumarin, ginsenoside R_{b1} , cryptotanshinone, tanshinone IIA, and oleanolic acid were purchased from the National Institute for the Control of Pharmaceutical and Biological Products (Beijing, P. R. China). Acetonitrile was of HPLC grade (Merck, Darmstadt, Germany). HPLC grade methanol was provided by Honeywell International Inc. (Burdick and Jackson, Muskegon, MI, USA). Phosphoric acid and acetic acid glacial were of HPLC grade and purchased from TianJin Chemical Reagents Development Center (TianJin, China). Ultrapure water for the preparation of samples and mobile phase was prepared with PURELAB Ultra GE MK2 water system (ELGA, High Wycombe, UK). Other reagents were of analytical grade. FTZ capsules were prepared by the Institute of Materia Medica, Guangdong

Pharmaceutical University (batch number 20090607). Eight comprised crude herbs were purchased from Zhixin Chinese Herbal Medicine Co., Ltd. (Guangzhou, China) and all the herbs were authenticated by Professor Shu-Yuan Li (Guangdong Pharmaceutical University). A voucher specimen was deposited in the Institute of Traditional Chinese Medicine, Guangdong Pharmaceutical University, Guangzhou, P. R. China.

Instrumentation and Analytical Conditions

The Waters AcQuityTM Ultra Performance LC system (Waters Corporation, Milford, USA) was equipped with quaternary pump, vacuum degasser, a cooling autosampler, and a diode-array detector. A UPLCTM BEH C18 column (50 × 2.1 mm, 1.7 μm) was utilized for separation with the column temperature at 30 °C. A binary gradient elution was adopted with mobile phase consisting of (A) 0.25% acetic acid glacial and 10 mM ammonium acetate in water and (B) acetonitrile: 0–1.6 min, B 2–5%; 1.6–7.6 min, B 5–20%; 7.6–9.6 min, B 20%; 9.6–14.6 min, B 20–35%; 14.6–17.6 min, B 35–80%; 17.6–18 min, B 80–100%; 18–18.4 min, B 100%. The flow rate was set at

Table 3 Structures of the Fructus Ligustri Lucidi constituents identified in FTZ preparation and serum samples from FTZ-treated rats

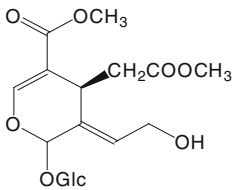
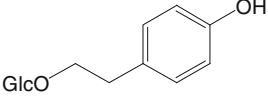
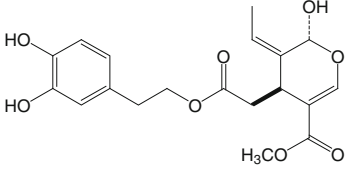
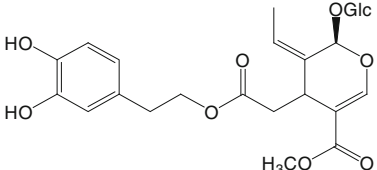
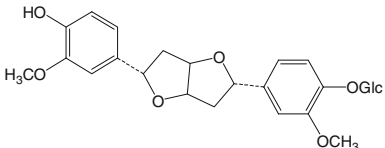
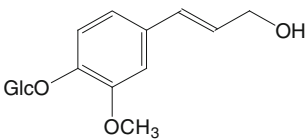
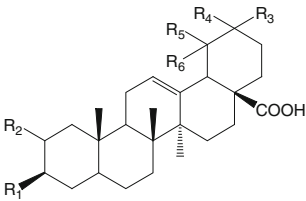
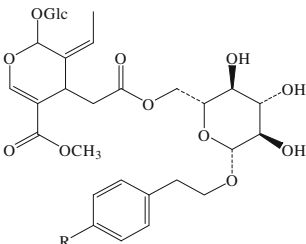
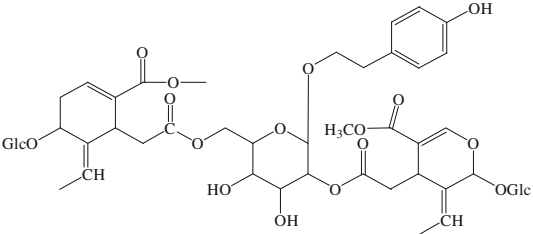
Type	Substituent group	Name of constituent	MW	Observed
		10-Hydroxyoleoside dimethyl ester	434	In serum and in preparation
		Salidroside	300	In serum and in preparation
		Oleuropein aglycone	378	In preparation only
		Oleuropein	540	In preparation only
		(+)-Pinoresinol-O- β -D-glucopyranoside	520	In preparation only
		Coniferin	342	In preparation only
	R ₁ =OH, R ₂ =OH, R ₃ =R ₄ =CH ₃ , R ₅ =R ₆ =H	Masilinic acid	472	In serum and in preparation
	R ₁ =CH ₃ COO, R ₂ =H, R ₃ =CH ₃ , R ₄ =H, R ₅ =CH ₃ , R ₆ =OH	Pomolic acid acetate	514	In serum and in preparation
	R ₁ =OH, R ₂ =H, R ₃ =R ₄ =CH ₃ , R ₅ =R ₆ =H	Oleanolic acid	456	In serum and in preparation
	R=OH	Specnuezhenide	686	In preparation only

Table 3 continued

Type	Substituent group	Name of constituent	MW	Observed
		Oleonuezhenide	1072	In preparation only

In serum and in preparation: the constituent was observed both in FTZ-treated serum and FTZ preparation, in preparation only: the constituent was only observed in FTZ preparation, in serum only: the constituent was only observed in FTZ-treated serum

MW molecular weight

0.40 mL min⁻¹. The autosampler was conditioned at 4 °C, and the injection volume was 10 µL.

The instrument Waters Micromass Q-TOF-microTM (Waters Co., UK) was equipped with the Lock Spray and ESI interface operating in both positive ion mode and negative ion mode, and with MassLynx data analysis software. The capillary voltage was set at 3 kV; the cone voltage was set at 30 V for both positive ionization mode and negative ionization mode. The ion source temperature was set at 100 °C and desolvation temperature at 350 °C. Nitrogen and argon were used for cone and collision gases, respectively. The cone and desolvation gas flows were 60 and 600 L h⁻¹, respectively. The mass spectrometric data was collected in full scan mode with the mass range of *m/z* 100–1,500, using independent reference lock-mass ions via the Lock Spray interface to ensure mass accuracy and reproducibility. The solution of chloramphenicol was used as lock-mass, with an [M+H]⁺ ion of *m/z* 345.0021 and an [M-H]⁻ ion of *m/z* 321.0045. The MS/MS analysis was performed using a variable collision energy (20–50 eV), which was optimized for each individual constituent. The Lock Spray frequency was set at 10 s. Acquity UPLC/Q-TOF micro system was operated using MassLynx 4.1 software (Waters Co., USA). The accurate mass and composition for the precursor and fragment ions were calculated by MassLynx 4.1.

Animals

Ten male Sprague–Dawley rats (body weight 200 ± 20 g) were obtained from the Medical Experimental Animal Center of Guangdong Province (Foshan, China). Animals were housed under standard conditions of temperature, humidity and light with food (laboratory rodent chow) and water provided ad libitum and were acclimated in the laboratory for at least 1 week prior to experiment. Before

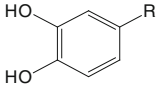
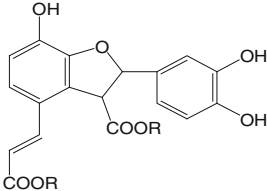
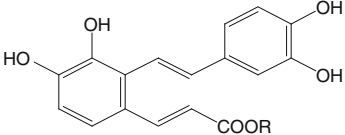
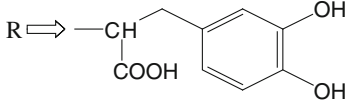
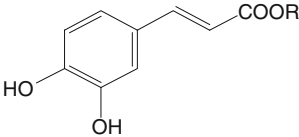
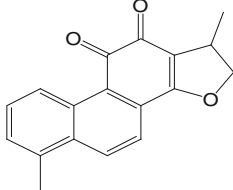
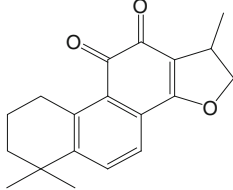
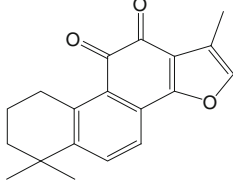
administration, the animals were fasted overnight with free access of water. All experimental protocols have been approved by Institutional Animal Ethics Committee of Guangdong Pharmaceutical University (GDPUIAEC No.200902), and are also in a compliance with national and international guidelines of animal welfare (NIH Guide for the Care and Use of Laboratory Animals, NIH publication No. 85-23, 1985).

Sample Preparation

Preparation of FTZ Extract

The preparation of FTZ extract from eight constituent herbs was consistent with the protocol described previously [1], and as follows: Radix Salvia Miltiorrhiza (50 g), Radix Atractylodes Macrocephala (60 g), Fructus Citri Sarcodactylis (50 g), Cortex Eucommiae (40 g), and Herba Cirsii Japonici (30 g) were extracted with boiling water twice (volume over weight values per reflux extraction = 12- and 9-fold, respectively; duration = 1.5 h per reflux extraction); Fructus Ligustri Lucidi (60 g) and Rhizoma Coptidis (20 g) were extracted with 70% ethanol twice (volume over weight values per reflux extraction = 10- and 8-fold, respectively; duration = 2 h per reflux extraction); Radix Notoginseng (20 g) was extracted with 50% ethanol twice (volume over weight values per reflux extraction = 10- and 8-fold, respectively; duration = 2 h per reflux extraction). The above three extracts were combined, filtered by gauzes, and the combined solution was freeze-dried. Five hundred milligrams of the freeze-dried powder was extracted with 50 mL methanol for 20 min under ultrasonics. The methanol extraction was centrifuged at 15,000 rpm for 15 min at 4 °C, and the supernatant was filtered through a 0.20-µm filter, the filtrate was applied for UPLC analysis. All authentic

Table 4 Structures of the Radix Salvia Miltiorrhiza constituents identified in FTZ preparation and serum samples from FTZ-treated rats

Type	Substituent group	Name of constituent	MW	Observed
	R=CH ₂ CH(OH)COOH	Danshensu	198	In preparation only
	R=CHO	Protocatechuic aldehyde	138	In preparation only
	R=COOH	Protocatechuic acid	154	In serum and in preparation
		Salvanolic acid B	718	In preparation only
	R ⇒ 	Salvanolic acid A	494	In preparation only
		Rosmarinic acid	360	In preparation only
		Dihydrotanshinone I	278	In preparation only
		Cryptotanshinone	296	In preparation only
		Tanshinone IIA	294	In preparation only

In serum and in preparation: the constituent was observed both in FTZ-treated serum and FTZ preparation, in preparation only: the constituent was only observed in FTZ preparation, in serum only: the constituent was only observed in FTZ-treated serum

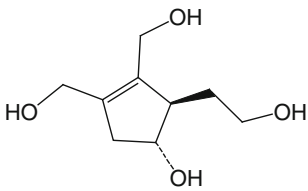
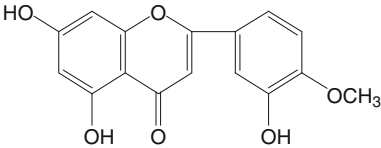
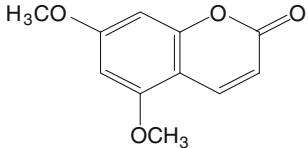
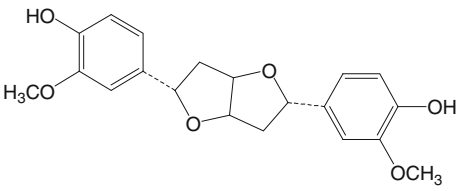
MW molecular weight

standards were accurately weighed, and dissolved in methanol to obtain stock solutions with indicated concentrations. All the stock solutions were stored in the refrigerator at 4 °C until analysis.

Preparation of Serum Samples

Capsule contents of FTZ, originated from the above extraction, were dispersed with distilled water as stock

Table 5 Structures of the remaining herbs constituents identified in FTZ preparation and serum samples from FTZ-treated rats

Type	Origin	Name of constituent	MW	Observed
	Cortex Eucommiae	Eucommiol	188	In serum and in preparation
	Herbal Cirsii Japonici and Fructus Citri Sarcodactylis	Diosmetin	300	In preparation only
	Fructus Citri Sarcodactylis	5,7-Dimethoxycoumarin	206	In serum and in preparation
	Cortex Eucommiae	Pinoresinol	358	In serum and in preparation

In serum and in preparation: the constituent was observed both in FTZ-treated serum and FTZ preparation; in preparation only: the constituent was only observed in FTZ preparation; in serum only: the constituent was only observed in FTZ-treated serum

MW molecular weight

solution (0.8 g mL^{-1}). The above suspension was orally administered to five rats ($2.0 \text{ mL}/100 \text{ g}$ body weight). An equal volume of distilled water was orally administered to the other five rats as control; 30 min after drug administration, the animals were anaesthetized by ether inhalation. The blood was collected from the vena ophthalmica and then centrifuged at 10,000 rpm for 5 min at $4 \text{ }^\circ\text{C}$. The supernatant obtained was frozen immediately and stored at $-80 \text{ }^\circ\text{C}$ before use.

Phosphoric acid ($240 \text{ } \mu\text{L}$) was added to 6.0 mL of the above supernatant and ultrasonicated for 1 min, and vortexed for 1 min. The mixed solution was applied to three pre-activated OASIS HLB solid phase extraction C18 columns (1 cc , 30 mg , Waters Corporation, USA). The column was washed with 4 mL of water, 2 mL of 100% methanol and 2 mL of 2% acetic acid glacial-methanol (1:9). The 100% methanol elutes and 2% acetic acid glacial-methanol (1:9) elutes were collected and dried under nitrogen gas at $50 \text{ }^\circ\text{C}$. The residues were re-dissolved in $300 \text{ } \mu\text{L}$ of methanol, centrifuged at 15,000 rpm for 15 min and an aliquot of supernatant was subjected to UPLC analysis.

Results and Discussions

UPLC-MS/MS Analysis and Identification of the Constituents of FTZ

ESI in both negative and positive ion modes was applied to analyze and identify the constituents in the FTZ. The total ion current chromatograms at the two ESI modes are shown in Fig. 1. Fifty-one peaks in FTZ were detected using UPLC-MS/MS, and 44 constituents were identified by comparing their retention behavior, the MS fragments characteristics to those of authentic standards. The names and structures of the identified constituents from Rhizoma Coptidis, Radix Notoginseng, Fructus Ligustri Lucidi, Radix Salvia miltiorrhiza, and other three herbs in both herbal preparation and the serum samples for FTZ-treated rats are listed in Tables 1, 2, 3, 4 and 5. The identified compounds are summarized in Table 6.

In order to obtain MS fragmentation patterns of constituents in FTZ, MS² spectra of 19 authentic standards were recorded by UPLC-MS/MS. Peaks 1, 3, 4, 6, 9, 10, 13, 14, 18, 20, 22, 23, 24, 25, 31, 33, 45, 49 and 51 were

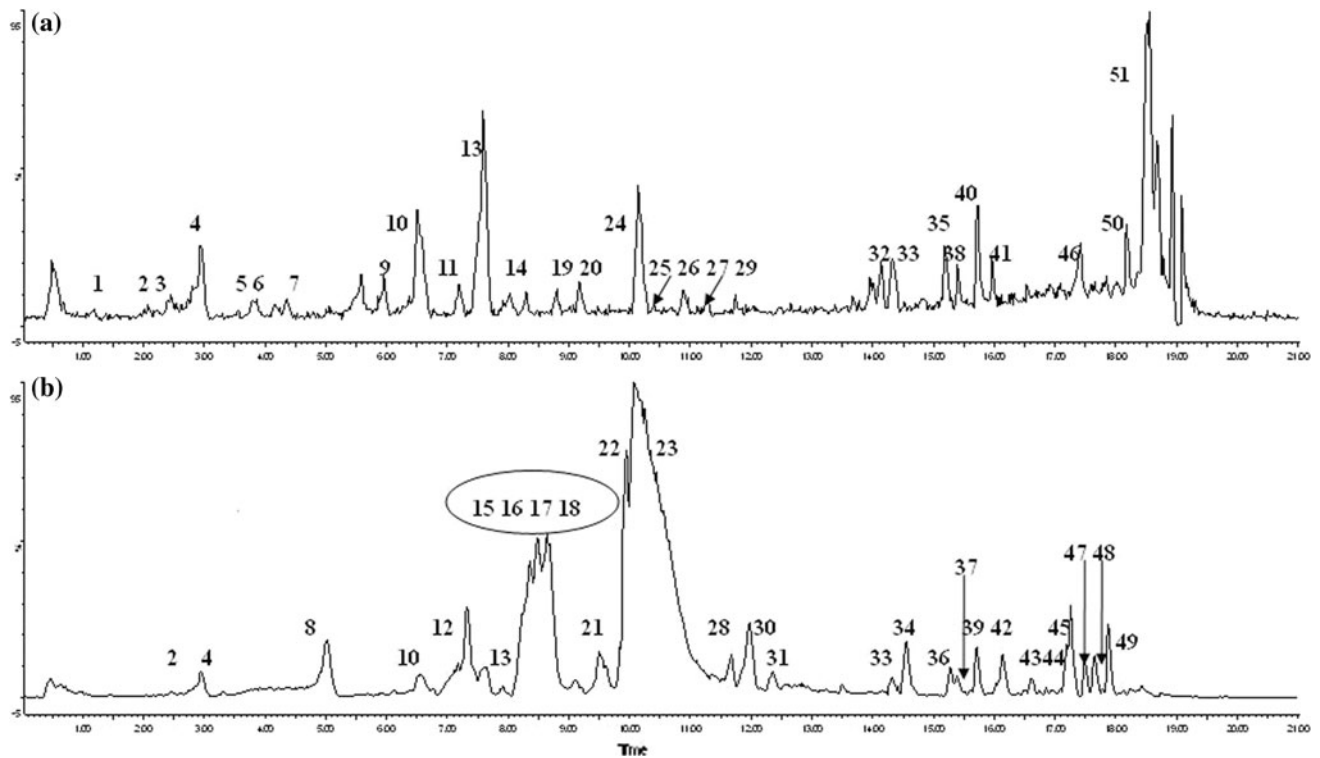


Fig. 1 UPLC-ESI-MS total ion current chromatograms of FTZ at the negative ion (a) and positive (b) mode. The numbers appearing in the circle. These peaks in the circle have similar structures and

retention times, each peak separate incompletely and cannot be marked clearly, the number in the circle represents four peaks including

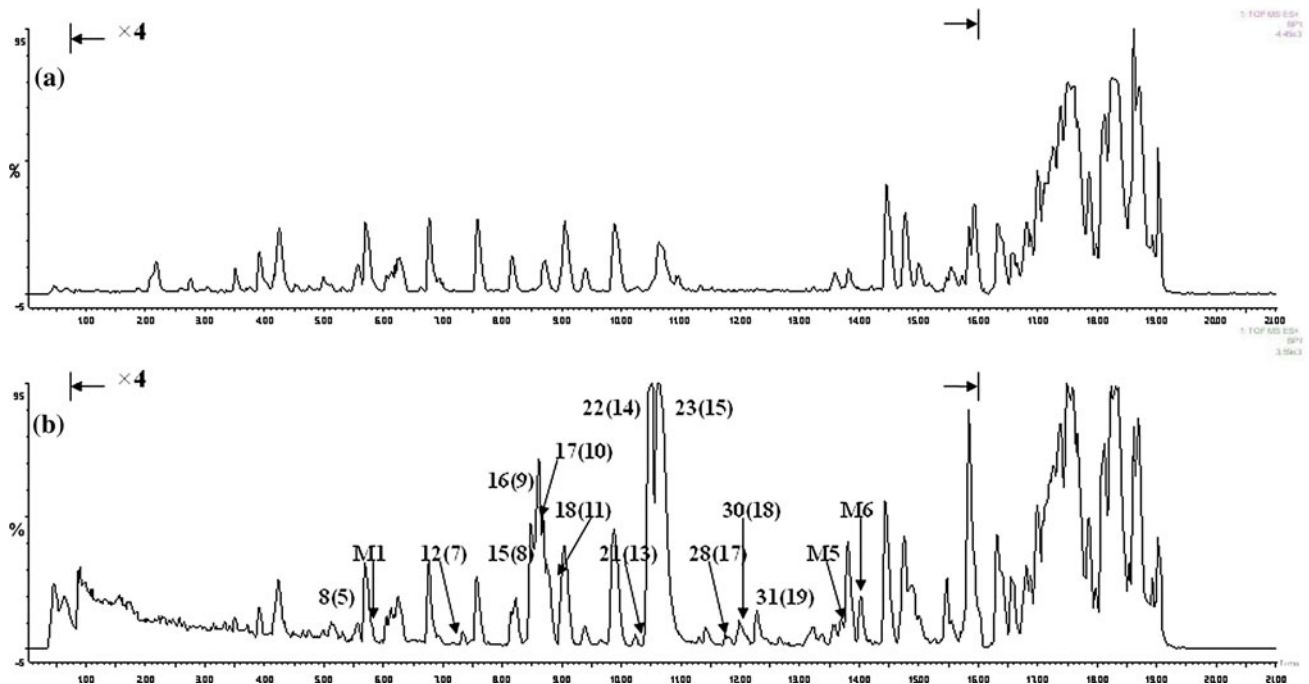


Fig. 2 Total ion current chromatograms at the positive ion mode of rat serum samples collected from control (a) and FTZ-treated (b) group

attributed to danshensu, protocatechuic acid, protocatechuic aldehyde, salidroside, rosmarinic acid, salvianolic acid B, specnuezhenide, salvianolic acid A,

jatrorrhizine, notoginsenoside R₁, palmatine, berberine, ginsenoside Rg₁, ginsenoside Re, 5,7-dimethoxycoumarin, ginsenoside Rb₁, cryptotanshinone, tanshinone IIA and

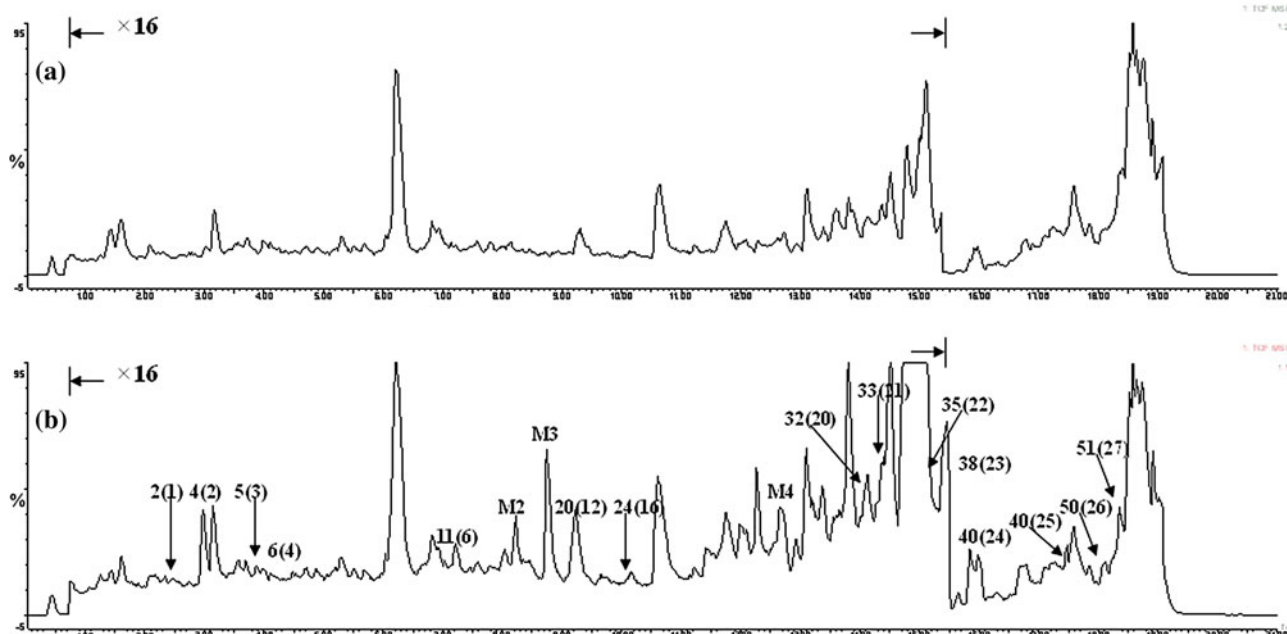


Fig. 3 Total ion current chromatograms at the negative ion mode of rat serum samples collected from control (a) and FTZ-treated (b) group

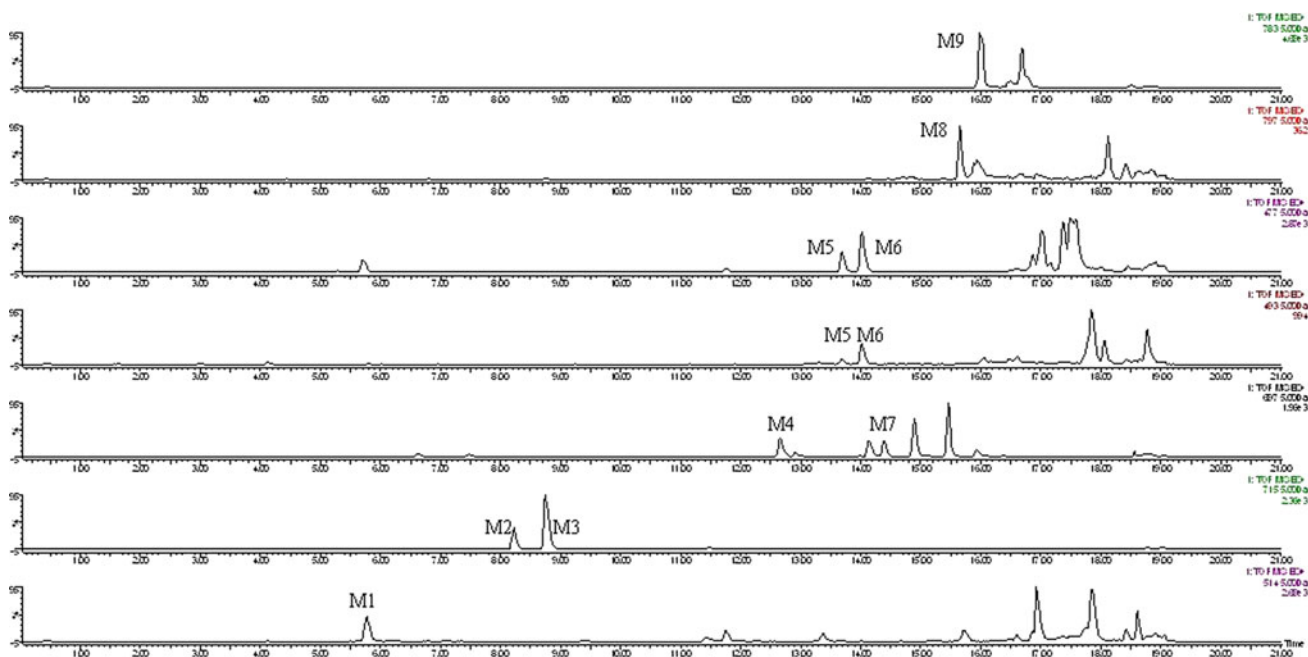


Fig. 4 Total ion current chromatograms of 9 metabolites identified from FTZ-treated rat serum samples

oleanolic acid, respectively, by comparing the retention time and mass data with those of the authentic standards. Other peaks were identified, utilizing elemental composition analysis of their MS and MS² data with software MassLynx from data and comparing with the literature data as well.

In the negative ion mode, ginsenosides, iridoid/secoiridoid glycosides, triterpene acids, and phenolic acids were observed in the FTZ, which originated from *Radix Notoginseng*, *Fructus Ligustri Lucidi* and *Radix Salvia*

miltiorrhiza, respectively. Among them, six ginsenosides, peaks 20, 24, 25, 32, 33, and 38, were identified as notoginsenoside R₁, ginsenoside Rg₁, ginsenoside Re, ginsenoside Rh₁/F₁ and ginsenoside Rb₁ and ginsenoside Rd, respectively, by comparison with authentic standards and literature data [17, 18]. The mass spectra of the ginsenosides exhibited the molecular ion peaks at [M-H]⁻ and [M-H+CH₃COOH]⁻. In the MS² spectra, aglycone ions *m/z* 475 and 459 were finally formed by loss of several

Table 6 MS data of (+) ESI-MS spectra and (-) ESI-MS spectra, and the identification results of the constituents of FTZ preparation and FTZ-treated rats serum

No.	Name	R_t (min)	MW	MS+ (m/z)	MS- (m/z)
1	Danshensu ^a	1.14	198		197[M-H] ⁻ 179[M-H-H ₂ O] ⁻ 135[M-H-H ₂ O-CO ₂] ⁻
2(1) ^b	10-Hydroxyoleoside dimethyl ester	2.45	434	452[M+NH ₄] ⁺	433[M-H] ⁻ 209[M-H-Glu-CH ₃ CO-H ₂ O] ⁻ 177[M-H-Glu-CH ₃ CO-CH ₃ O-H ₂ O] ⁻ 165[M-H-Glu-2CH ₃ CO-H ₂ O] ⁻
3	Protocatechuic aldehyde ^a	2.62	138		137[M-H] ⁻ 109[M-H-CO] ⁻
4(2)	Salidroside ^a	2.99	300	318[M+NH ₄] ⁺	299[M-H] ⁻ 179[M-H-C ₈ H ₈ O] ⁻ 119[M-H-C ₆ H ₁₂ O ₆] ⁻ 101[M-H-Glu] ⁻
5(3)	Hydroxyl-palmitate	3.85	368		367[M-H] ⁻ 352; 191
6(4)	Protocatechuic acid ^a	3.89	154		153[M-H] ⁻ 109[M-H-CO ₂] ⁻
7	Oleuropein aglycone	4.36	378		377[M-H] ⁻ 197[M-H-C ₉ H ₈ O ₄] ⁻ ; 153
8(5)	Magnoflorine	5.04	342	342[M] ⁺ 297[M+H-CH ₃ -CH ₃ O] ⁺ 282[M+H-2CH ₃ -CH ₃ O] ⁺ 265[M-CH ₃ -2CH ₃ O] ⁺ 237[M-CH ₃ -2CH ₃ O-H ₂ O] ⁺	
M1	Jatrorrhizine 3-O-β-D-glucuronide	5.77	514	514[M] ⁺ 338[M-GlcUA] ⁺	
9	Rosmarinic acid ^a	5.96	360		359[M-H] ⁻ 197[M-H-C ₉ H ₆ O ₃] ⁻ 179[M-H-C ₉ H ₆ O ₃ -H ₂ O] ⁻ 161[M-H-C ₉ H ₆ O ₃ -2H ₂ O] ⁻
10	Salvianolic acid B ^a	6.52	718	736[M+NH ₄] ⁺	717[M-H] ⁻ 519[M-H-C ₉ H ₁₀ O ₅] ⁻ 321[M-H-2C ₉ H ₁₀ O ₅] ⁻ 339[M-H-C ₉ H ₁₀ O ₅ -C ₉ H ₈ O ₄] ⁻
11(6)	Eucommiol	7.19	188		187[M-H] ⁻ ; 125; 97
12(7)	Thalifendine	7.33	322	322[M] ⁺ 307[M-CH ₃] ⁺ 279[M-CO-CH ₃] ⁺ 251[M-2CO-CH ₃] ⁺	
13	Specnuezhenide ^a	7.59	686	704[M+NH ₄] ⁺	685[M-H] ⁻ 523[M-H-Glu] ⁻ 453[M-H-Glu-C ₃ H ₂ O ₂] ⁻ 423[M-H-Glu-C ₃ H ₂ O ₂ -CH ₂ O] ⁻ 299[M-H-Glu-C ₃ H ₂ O ₂ -CH ₂ O-C ₇ H ₈ O ₂] ⁻ ; 223; 197

Table 6 continued

No.	Name	R_t (min)	MW	MS+ (m/z)	MS- (m/z)
14	Salvianolic acid A ^a	8.03	494		493[M-H] ⁻ 313[M-H-C ₉ H ₈ O ₄] ⁻ 295[M-H-C ₉ H ₁₀ O ₅] ⁻ 185[M-H-C ₉ H ₁₀ O ₅ -C ₆ H ₆ O ₂] ⁻
M2	25-Hydroxy-ginsenoside Rh ₁ /F ₁	8.23	656		715[M-H+CH ₃ COOH] ⁻ 655[M-H] ⁻ ; 493[M-H-Glu] ⁻
15(8)	Columbamine	8.33	338	338[M] ⁺ 322[M-CH ₄] ⁺ 308[M-2CH ₃] ⁺ 294[M-CH ₄ -CO] ⁺ 280[M-2CH ₃ -CO] ⁺	
16(9)	Epiberberine	8.41	336	336[M] ⁺ 320[M-CH ₄] ⁺ 294[M-CH ₄ -CO] ⁺ 263[M-CH ₃ -CH ₂ O-CO] ⁺	
17(10)	Coptisine	8.47	320	320[M] ⁺ 292[M-CO] ⁺ 262[M-CO-CH ₂ O] ⁺ 234[M-2CO-CH ₂ O] ⁺ 204[M-2CO-2CH ₂ O] ⁺	
18(11)	Jatrorrhizine ^a	8.66	338	338[M] ⁺ 322[M-CH ₄] ⁺ 308[M-2CH ₃] ⁺ 294[M-CH ₄ -CO] ⁺ 280[M-2CH ₃ -CO] ⁺ 265[M-CH ₃ -CO-CH ₂ O] ⁺ 250[M-2CH ₃ -CO-CH ₂ O] ⁺	
M3	25-Hydroxy-ginsenoside Rh ₁ /F ₁	8.74	656		715[M-H+CH ₃ COOH] ⁻ 655[M-H] ⁻ ; 493[M-H-Glu] ⁻
19	Oleuropein	8.81	540		539[M-H] ⁻ 377[M-H-Glu] ⁻ 307[M-H-Glu-C ₄ H ₆ O] ⁻ 275[M-H-Glu-C ₄ H ₆ O-CH ₃ OH] ⁻
20(12)	Notoginsenoside R ₁ ^a	9.17	932		991[M-H+CH ₃ COOH] ⁻ 931[M-H] ⁻ 799[M-H-Xyl] ⁻ 637[M-H-Xyl-Glu] ⁻ 475[M-H-Xyl-2Glu] ⁻
21(13)	Berberrubine	9.49	322	322[M] ⁺ 307[M-CH ₃] ⁺ 279[M-CO-CH ₃] ⁺ 251[M-2CO-CH ₃] ⁺	

Table 6 continued

No.	Name	R_t (min)	MW	MS+ (m/z)	MS- (m/z)
22(14)	Palmatine ^a	9.93	352	352[M] ⁺ 337[M-CH ₃] ⁺ 336[M-CH ₄] ⁺ 322[M-2CH ₃] ⁺ 308[M-CH ₄ -CO] ⁺ 294[M-2CH ₃ -CO] ⁺ 279[M-CH ₃ -CO-CH ₂ O] ⁺	
23(15)	Berberine ^a	10.06	336	336[M] ⁺ 321[M-CH ₃] ⁺ 320[M-CH ₄] ⁺ 306[M-2CH ₃] ⁺ 292[M-CH ₄ -CO] ⁺ 278[M-2CH ₃ -CO] ⁺ 262[M-CH ₄ -CO-CH ₂ O] ⁺ 234[M-CH ₄ -2CO-CH ₂ O] ⁺	
24(16)	Ginsenoside Rg ₁ ^a	10.14	800		859[M-H+CH ₃ COOH] ⁻ 637[M-H-Glu] ⁻ ; 475[Ag] ⁻
25	Ginsenoside Re ^a	10.39	946		1005[M-H+CH ₃ COOH] ⁻ 945[M-H] ⁻ 799[M-H-Rha] ⁻ 637[M-H-Rha-Glu] ⁻ 475[M-H-Rha-2Glu] ⁻
26	Diosmetin	10.88	300		299[M-H] ⁻ 284[M-H-CH ₃] ⁻ 256[M-H-CH ₃ -CO] ⁻
27	(+)-Pinoresinol-O-β-D-glucopyranoside	11.27	520	521[M+H] ⁺	519[M-H] ⁻ 357[M-H-Glu] ⁻ 235[M-H-Glu-CH ₂ O-C ₆ H ₄ O] ⁻
28(17)	Dehydrocorydaline	11.67	366	366[M] ⁺ 350[M-CH ₄] ⁺ 336[M-2CH ₃] ⁺ 322[M-CH ₄ -CO] ⁺ 308[M-2CH ₃ -CO] ⁺ 278[M-2CH ₃ -CO-CH ₂ O] ⁺	
29	Oleonuezhenide	11.74	1072		1071[M-H] ⁻ 771[M-H-C ₁₄ H ₂₀ O ₇] ⁻ 685[M-H-C ₁₇ H ₂₂ O ₁₀] ⁻ 523[M-H-C ₁₇ H ₂₂ O ₁₀ -Glu] ⁻
30(18)	13-Methylberberine	11.97	350	350[M] ⁺ 335[M-CH ₃] ⁺ 334[M-CH ₄] ⁺ 320[M-2CH ₃] ⁺ 306[M-CH ₄ -CO] ⁺ 292[M-2CH ₃ -CO] ⁺ 278[M-CH ₄ -2CO] ⁺	

Table 6 continued

No.	Name	R_t (min)	MW	MS+ (m/z)	MS- (m/z)
31(19)	5,7-Dimethoxycoumarin ^a	12.35	206	207[M+H] ⁺ 192[M+H-CH ₃] ⁺ 164[M+H-CH ₃ -CO] ⁺ 149[M+H-2CH ₃ -CO] ⁺	
M4	20(S)-Ginsenoside Rh ₁ / 20(R)-Ginsenoside Rh ₁ / Ginsenoside F ₁	12.66	638		697[M-H+CH ₃ COOH] ⁻ 441; 423; 405
M5	20(S)/(R)-Protopanaxatriol	13.69	476	477[M+H] ⁺	493[M-H+H ₂ O] ⁻ 553[M-H+H ₂ O+CH ₃ COOH] ⁻
M6	20(S)/(R)-Protopanaxatriol	14.02	476	477[M+H] ⁺	493[M-H+H ₂ O] ⁻ 553[M-H+H ₂ O+CH ₃ COOH] ⁻
32(20)	Ginsenoside F ₁ / Ginsenoside Rh ₁	14.14	638		697[M-H+CH ₃ COOH] ⁻ 637[M-H] ⁻ 475[M-H-Glu] ⁻
33(21)	Ginsenoside Rb ₁ ^a	14.31	1108	1109[M+H] ⁺ 1126[M+NH ₄] ⁺ 1091[M+H-H ₂ O] ⁺	1107[M-H] ⁻ 1167[M-H+CH ₃ COOH] ⁻ 945[M-H-Glu] ⁻ 783[M-H-2Glu] ⁻ 603[M-H-3Glu-H ₂ O] ⁻ ; 459[Ag] ⁻
M7	20(S)-Ginsenoside Rh ₁ / 20(R)-Ginsenoside Rh ₁ / Ginsenoside F ₁	14.39	638		697[M-H+CH ₃ COOH] ⁻ 441; 423; 405
34	Coniferin	14.52	342	343[M+H] ⁺ ; 295 181[M+H-Glu] ⁺ 164[M+H-Glu-OH] ⁺ ; 120	
35(22)	Not identified	15.18	532		531[M-H] ⁻ 1063[2M-H] ⁻ 489[M-H-CH ₂ -CO] ⁻ 471[M-H-CH ₂ -CO-H ₂ O] ⁻ 427[M-H-CH ₂ -CO-H ₂ O-CO ₂] ⁻
36	Not identified	15.27		357; 328; 293; 249	
37	13-Ethyl-5,6-dihydro-2,3,9,10-tetramethoxy-dibenzo[a,g]quinolizinium	15.36	380	398[M+NH ₄] ⁺ 366[M-CH ₂] ⁺ 351[M-CH ₂ -CH ₃] ⁺ 336[M-CH ₂ -2CH ₃] ⁺ 322[M-CH ₂ -CH ₄ -CO] ⁺	
38(23)	Ginsenoside Rd	15.39	946	947[M+H] ⁺ ; 965[M+NH ₄] ⁺	945[M-H] ⁻ 1005[M-H+CH ₃ COOH] ⁻ 783[M-H-Glu] ⁻ 621[M-H-2Glu] ⁻ ; 459[Ag] ⁻
M8	Salvianolic acid B sulfates	15.65	798		797[M-H] ⁻ 717[M-H-SO ₃] ⁻

Table 6 continued

No.	Name	R_t (min)	MW	MS+ (m/z)	MS- (m/z)
39	Dehydrocorybulbine	15.70	352	352[M] ⁺ 337[M-CH ₃] ⁺ 322[M-2CH ₃] ⁺ 308[M-CH ₄ -CO] ⁺ 294[M-2CH ₃ -CO] ⁺	
40(24)	Pinoresinol	15.71	358		357[M-H] ⁻ 342[M-H-CH ₃] ⁻ 311[M-H-CH ₃ -2CH ₃ O] ⁻ ; 151
41	Not identified	15.96			259; 244; 201; 189
M9	Ginsenoside Rg ₃	15.97	784		783[M-H] ⁻ 621[M-H-Glu] ⁻ 459[M-H-2Glu] ⁻
42	Not identified	16.14		614; 336; 321	
43	Dihydrotanshinone I	16.60	278	279[M+H] ⁺ 261[M+H-H ₂ O] ⁺ 233[M+H-H ₂ O-CO] ⁺	
44	Not identified	17.19	519	520[M+H] ⁺ 502[M+H-H ₂ O] ⁺	578[M-H+CH ₃ COOH] ⁻
45	Cryptotanshinone ^a	17.24	296	297[M+H] ⁺ 279[M+H-H ₂ O] ⁺ 268[M+H-C ₃ H ₆] ⁺ 251[M+H-H ₂ O-CO] ⁺ 227[M+H-C ₂ H ₅] ⁺	
46(25)	Masilinic acid	17.39	472		471[M-H] ⁻ 411[M-2H-CH ₃ -CO ₂] ⁻ 393[M-2H-CH ₃ -CO ₂ -H ₂ O] ⁻
47	Not identified	17.49		496; 478; 184; 104	
48	Not identified	17.64		522; 503; 184; 104	
49	Tanshinone IIA ^a	17.87	294	295[M+H] ⁺ 277[M+H-H ₂ O] ⁺ 249[M+H-H ₂ O-CO] ⁺ 266[M+H-C ₂ H ₅] ⁺	
50(26)	Pomolic acid acetate	18.19	514		513[M-H] ⁻ 495[M-H-H ₂ O] ⁻ 453[M-H-CH ₂ CO-H ₂ O] ⁻
51(27)	Oleanolic acid ^a	18.51	456		455[M-H] ⁻ 407[M-H-HCHO-H ₂ O] ⁻ 391[M-H-HCOOH-H ₂ O] ⁻ 373[M-H-HCOOH-2H ₂ O] ⁻

MW molecular weight, *Glu* glucose, *Rha* rhamnose, *Xyl* xylose, *Agl* aglycone, *GlcUA* glucuronic acid, *M* metabolites in serum after oral administration of FTZ

^a Structure confirmed by comparison with authentic standards

^b Number in parenthesis represents the no. of peak detected in serum after oral administration of FTZ

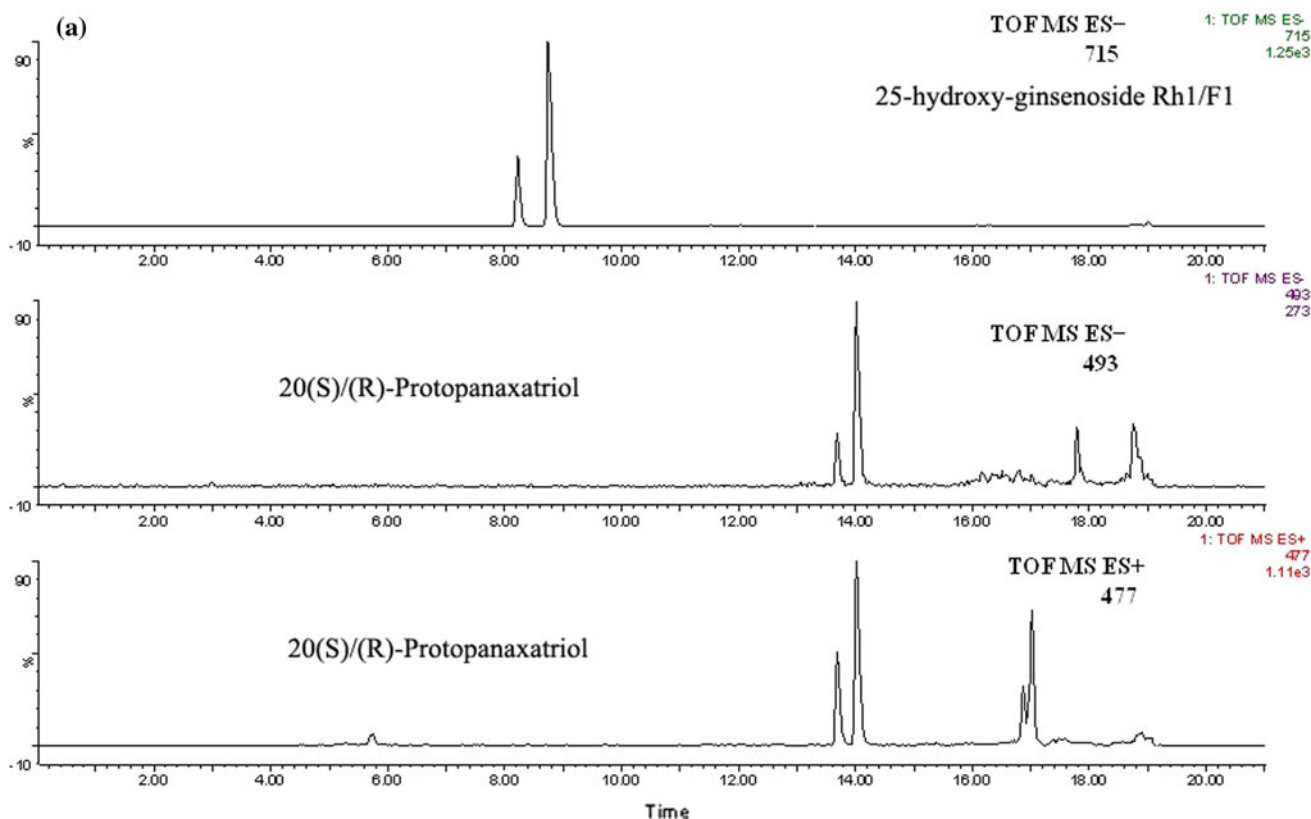


Fig. 5 EIC (a), MS (b) and MS² (c) spectra of 25-hydroxy-ginsenoside Rh₁/F₁ and protopanaxatriol and the possible metabolic biotransformation pathways (d)

glycosidic units, which were the characteristic ions of panaxatriols and panaxadiols, respectively [19]. Thus, these peaks could be identified as ginsenosides. For example, peak 24 showed a molecular ion at m/z 859 $[M-H+CH_3COOH]^-$ in MS spectra and exhibited m/z 637 $[M-H-Glu]^-$ and m/z 475 $[Ag]^-$ ions in the MS² spectra. The fragmentation ion at m/z 475 was produced by loss of all linked glucosidic bonds, which was a characteristic fragmentation of protopanaxatriol type ginsenosides [19]. Peak 33 showed a molecular ion at m/z 1107 $[M-H]^-$ in MS spectra; m/z 945 $[M-H-Glu]^-$, m/z 783 $[M-H-2Glu]^-$, m/z 603 $[M-H-3Glu-H_2O]^-$ and m/z 459 $[Ag]^-$ ions could be detected in the MS² spectra, which exhibited a fragmentation pathway corresponding to the loss of glycosidic units. The fragmentation ion at m/z 459 corresponds to a characteristic ion of the protopanaxadiol moiety [20].

Iridoid glycosides, secoiridoid glycosides and triterpene acids are the essential constituents in the Fructus Ligustri Lucidi extract of FTZ, which include salidroside, oleuropein aglycone, oleuropein, specnuezhenide, masilinic acid, pomolic acid acetate, oleanolic acid [21, 22]. Peak 13 showed a molecular ion at m/z 685 $[M-H]^-$ in MS spectra and exhibited m/z 523, 453, 423, 299, 223 and 197 ions in the MS² spectra. By comparison with the authentic

standard, peak 13 was unambiguously identified as specnuezhenide. The identification of peak 19 as oleuropein was corroborated by detection of the molecular ion at m/z 539 and its aglycone fragment at m/z 377. The MS spectrum showed a quasi-molecular ion at m/z 539 $[M-H]^-$ and the fragments were consistent with the following fragmentation pattern: the ion at m/z 377 arose from the loss of glucose, the ion at m/z 307 was characteristic of the loss of a C_4H_6O fragment and the fragment at m/z 275 might derive from the loss of CH_3OH from the elenolic fragment of the molecule [21]. Peak 7 exhibited the pseudo-molecular ion at m/z 377 in MS and characteristic ions at m/z 197 and m/z 153 in its MS² spectrum, corresponding to the oleuropein aglycone or its isomer. By retrieving of literature data [22], peak 7 was identified as oleuropein aglycone.

Among 51 analytes, there are six phenolic acids and three diterpenoids originated from Radix Salvia Miltiorrhiza. Phenolic acids could be classified into monomer and polymer. Polymers could be composed of one or several different monomers such as danshensu, caffeic acid or others. In the MS² spectra of three monomer standards, including small molecules such as CO_2 , CO and H_2O were produced in the fragmentation pathways, which indicated

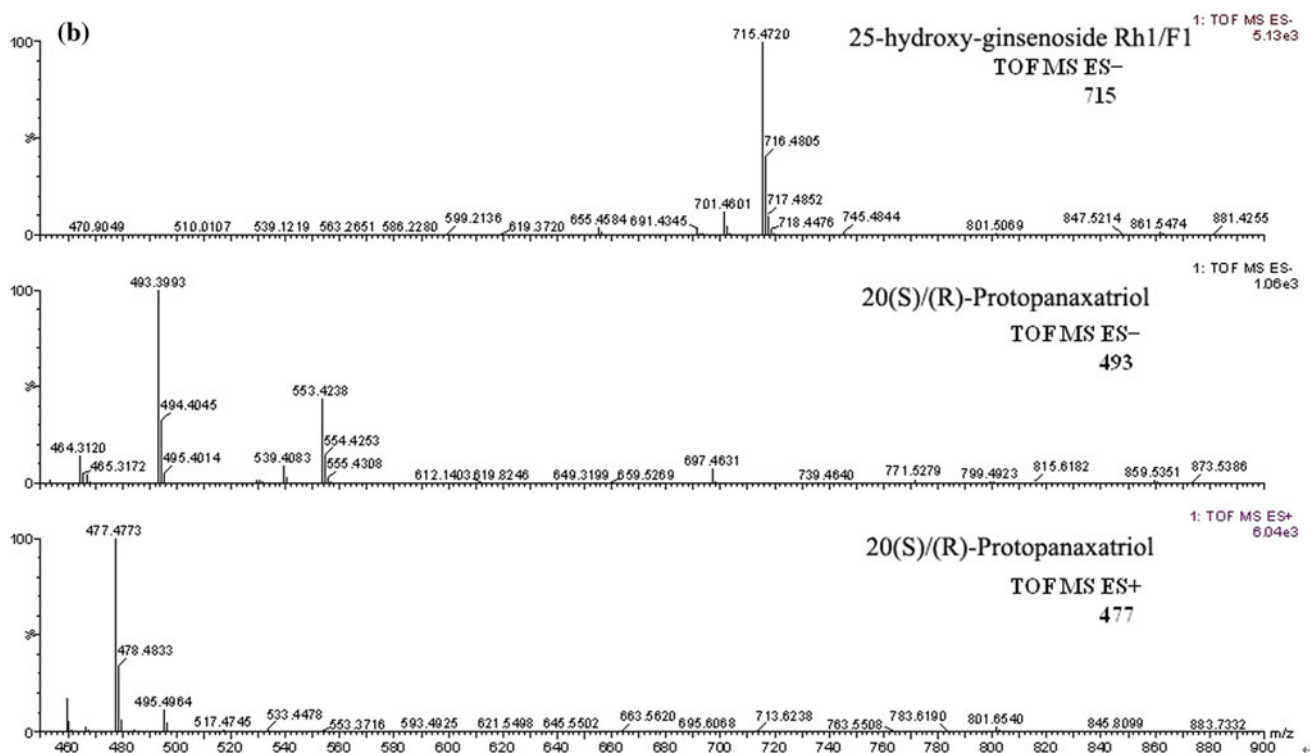


Fig. 5 continued

the presence of carboxyl, carbonyl or hydroxyl groups [23]. Danshensu showed a $[M-H]^-$ ion at m/z 197, and produced m/z 179 $[M-H-H_2O]^-$ and m/z 135 $[M-H-H_2O-CO_2]^-$. Similar to danshensu, both of the $[M-H]^-$ ions at m/z 137 of protocathechuic aldehyde and m/z 153 of protocathechuic acid produced the same ion at m/z 109 corresponding to the loss of CO and CO₂, respectively.

As to three polymers, which contained an ester bond or ester bonds, the predominant fragmentation of their $[M-H]^-$ ions was the cleavage of the ester bond to lose danshensu $[M-H-198]^-$ and caffeic acid $[M-H-180]^-$. For instance, peak 10 exhibited a quasi-molecular ion $[M-H]^-$ of m/z 717. Its MS² spectra gave rise to prominent ion at m/z 519 corresponding to the loss of a molecule of danshensu. Other two fragment ions, $[M-H-198-198]^-$ ion at m/z 321 and $[M-H-198-180]^-$ ion at m/z 339 corresponding to the loss of the second danshensu and the first caffeic acid. These data are consistent with those in the literature [23]. Therefore, peak 10 was tentatively identified as salvianolic acid B. Similarly, peaks 9, 14 were identified as rosmarinic acid and salvianolic acid A separately [23, 24].

Rhizoma Coptidis alkaloids, which were the most abundant constituents in the alcohol extra of FTZ, exhibited a special fragmentation pathway in the positive ion mode. It is well known that loss the neutral species such as CO, CH₃, CH₄ and CH₂O were observed in the MS² spectra of Rhizoma Coptidis alkaloids [9]. Peak 23 showed

a molecular ion at m/z 336 $[M]^+$ in MS spectra, and exhibited some ions at m/z 320 $[M-CH_4]^+$, 306 $[M-2CH_3]^+$, 292 $[M-CH_4-CO]^+$, 278 $[M-2CH_3-CO]^+$, 262 $[M-CH_4-CO-CH_2O]^+$ and 234 $[M-CH_4-2CO-CH_2O]^+$ in MS² spectra, showing the neutral loss of CO, CH₃, CH₄ and CH₂O in the fragmentation pathway. These data are typical for the Rhizoma Coptidis alkaloids in the present study and consistent with those in the literature [9]. Thus, the compound was identified as berberine. Similarly, peaks 8, 12, 15, 16, 17, 18, 21, 22, 28, 30 and 39 were identified as magnoflorine, thalifendine, columbamine, epiberberine, coptisine, jatrorrhizine, berberrubine, palmatine, dehydrocorydaline, 13-methylberberine and dehydrocorybulbine, respectively [9, 25–27]. Peak 37 showed the molecular ion at m/z 398 $[M+NH_4]^+$ and its product typical fragments at m/z 366 $[M-CH_2]^+$, 351 $[M-CH_2-CH_3]^+$, 336 $[M-CH_2-2CH_3]^+$ and 322 $[M-CH_2-CH_4-CO]^+$ respectively in the MS² spectrum. According to the literature data [28], we suggested that peak 37 could be 13-ethyl-5,6-dihydro-2,3,9,10-tetramethoxy-dibenzo[a,g]quinolizinium.

In addition to Rhizoma Coptidis alkaloids in positive ion mode, three diterpenoids (dihydrotanshinone I, cryptotanshinone and tanshinone IIA) also exhibited $[M+H]^+$ ions in positive ion mode. It is well known that hydrogen at C-1 and oxygen at C-11 of tanshinones were the source of the dissociated H₂O and the neutral species such as CO, H₂O,

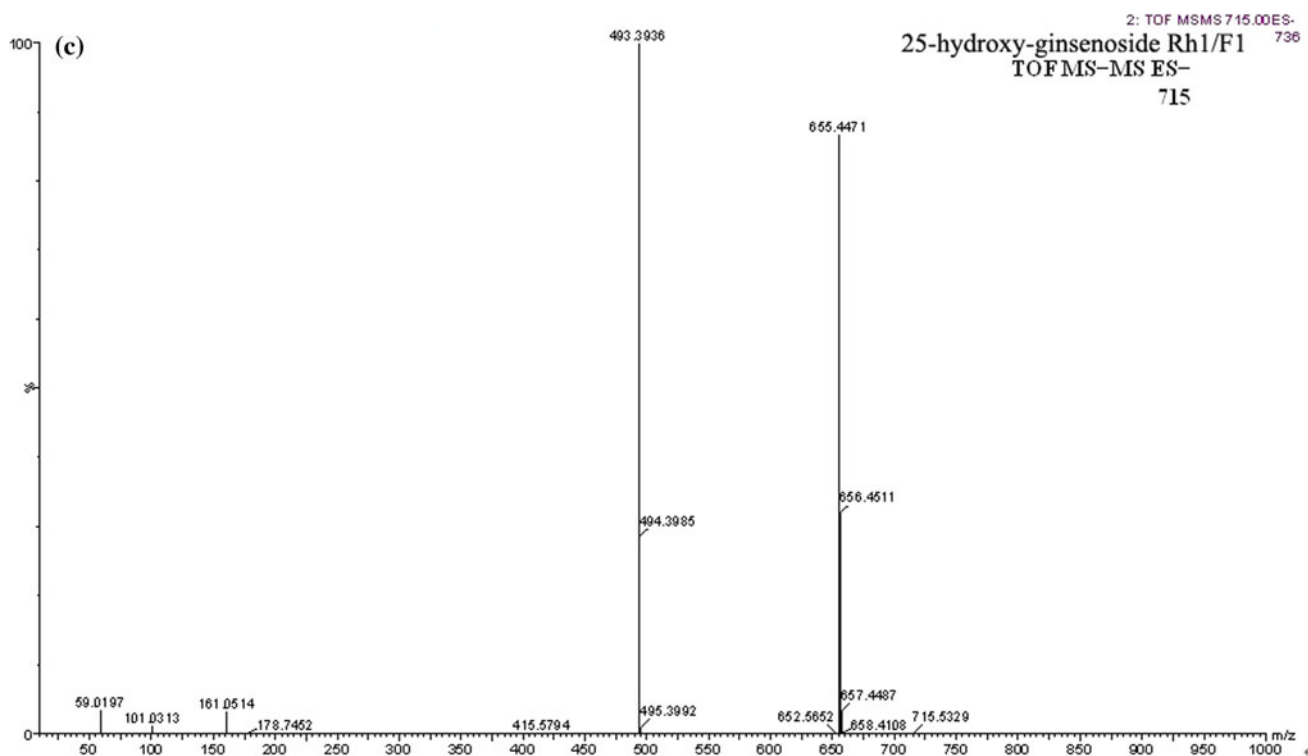


Fig. 5 continued

C_2H_5 and C_3H_6 were also observed in the MS^2 spectra [29]. Peak 45 showed a molecular ion at m/z 297 $[M+H]^+$ in MS spectra, and exhibited an ion at m/z 279 $[M+H-H_2O]^+$ in MS^2 spectra, which corresponded to three fragment ions at m/z 268 $[M+H-C_3H_6]^+$, m/z 227 $[M+H-C_2H_5]^+$ and m/z 251 $[M+H-H_2O-CO]^+$, showing the neutral loss of CO, H_2O , C_2H_5 and C_3H_6 in the fragmentation pathway. According to these data, peak 45 was tentatively identified as cryptotanshinone [30]. Using the same method, peak 43 and peak 49 were identified as dihydrotanshinone I and tanshinone IIA by comparison with literature data and authentic standards [30, 31].

In addition, the molecular ion $[M+H]^+$ of peak 31 was observed in the MS spectra, which dissociated in MS^2 to generate several ions at m/z 192, 164, 149 and 121. The ion at m/z 192 can be attributed to the loss of a methyl radical from the parent ion, this ion fragmented further with the loss of CO to give a signal at m/z 164. Subsequent loss of a methyl and a CO group radical to exhibited ions at m/z 149 and 121, were observed. Comparing with the authentic standard and literature data [32], peak 31 was tentatively identified as 5, 7-dimethoxycoumarin. Peak 34 showed a molecular ion at m/z 343 $[M+H]^+$ in MS spectra, and exhibited four ions at m/z 295, m/z 181 $[M+H-Glu]^+$, m/z 164 $[M+H-Glu-OH]^+$ and m/z 120 in MS^2 spectra, showing the loss of glucoside and hydroxy group in the fragmentation pathway. By

comparison with literature data [33], this component was ascertained as coniferin.

UPLC-MS/MS Analysis and Identification of the Constituents of FTZ in Rat Serum

Analysis of FTZ Constituents in Rat Serum

By comparison with the mass chromatography of FTZ and the rat serum samples from control group, the MS spectra for rat serum samples from FTZ-treated group exhibited 27 peaks in common, which demonstrated that the 27 components from FTZ were absorbed into the rat blood after oral administration. In addition, there were another nine peaks, which were only detected in the dosed serum, indicating that those components were metabolites of constituents from FTZ. Ion chromatograms of dosed and controlled rat serum are shown in Figs. 2, 3 and 4. The MS spectra and retention behavior of 36 peaks for prototype components and metabolites are summarized in Table 6.

Analysis of Prototype Constituents of FTZ in Rat Serum

The constituents in rat serum after oral administration of FTZ were identified using their retention time and mass spectra. As a result, peaks 1, 2, 22, 26 and 27 were original form compounds existing in *Fructus Ligustri Lucidi*; peaks

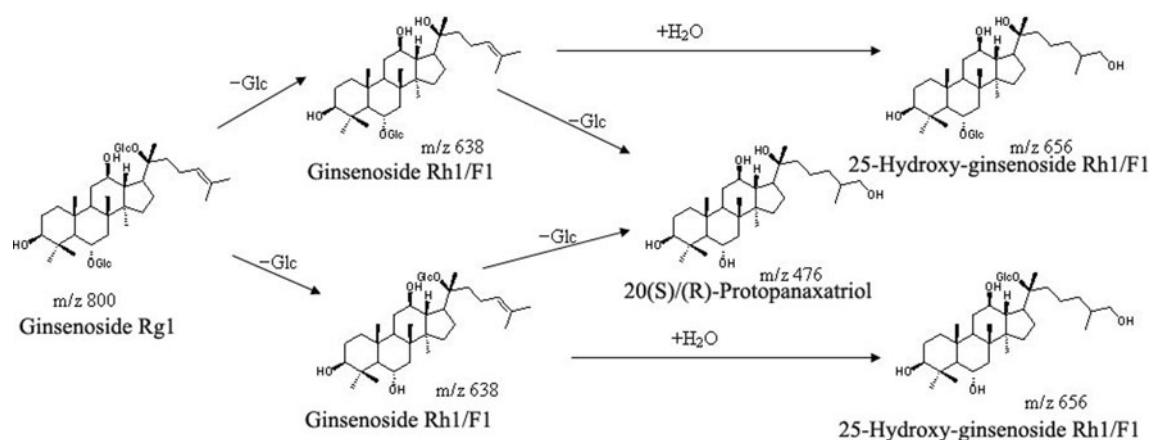


Fig. 5 continued

3, 5, 7, 8, 9, 10, 11, 13, 14, 15, 17 and 18 came from *Rhizoma Coptidis*; peaks 12, 16, 20, 21 and 23 resulted from *Radix Notoginseng*; peak 19 and 22 originated from *Fructus Citri Sarcodactylis*; peak 6 and 24 came from *Cortex Eucommiae*; peak 4 originated from *Radix Salvia Miltiorrhiza*. It displayed that most of alkaloids, ginsenosides and pentacyclic triterpenes could be unambiguously detected in their original forms from the rat serum after FTZ administration.

Analysis of Metabolites of FTZ in Rat Serum

To identify the metabolites accurately, probable structures were first postulated in accordance with the rules and characteristics of drug metabolism *in vivo*. In this study, the constituents of FTZ extract have been identified. These data may provide guidance for investigating the metabolites of FTZ in rat serum. M1 was identified as the glucuronide conjugate of alkaloids, jatrorrhizine-3-O- β -D-glucuronide, since it showed the m/z 514 [M]⁺ in MS spectra, and exhibited m/z 338 [M-GlcUA]⁺ in MS² spectra, which was confirmed by comparison with literature data [34]. M2 and M3 were suspected to be metabolite of ginsenoside Rh₁/F₁, both of them showed the same molecular ion at m/z 715 [M-H+CH₃COOH]⁻ in MS spectra, and exhibited product ions m/z 655 [M-H]⁻ and m/z 493 [M-H-Glu]⁻ in MS² spectra. By comparison with the literature data [35, 36], this showed the same fragmentation pathway as the metabolite of ginsenoside Rh₁/F₁, so the two constituents were identified as the 25-hydroxyl-ginsenoside Rh₁/F₁. Using the same method, M5 and M6 were identified as 20(S)/(R)-protopanaxatriol because they showed the m/z 477 [M+H]⁺ ion in positive ion mode and m/z 493 [M-H+H₂O]⁻ and m/z 553 [M-H+H₂O+CH₃COOH]⁻ ions in negative ion mode. By comparison with the literature data [36], we suggested that M5 and M6 may be sapogenin which formed

by loss of all glycosidic units from protopanaxatriol saponins. The MS and MS² spectra and possible metabolic pathways of 25-hydroxy-ginsenoside Rh₁/F₁ and protopanaxatriol in positive and negative ion mode are shown in Fig. 5a-d. M4 and M7 showed the molecular ion at m/z 697 [M-H+CH₃COOH]⁻ in MS spectra, and exhibited m/z 441, 423 and 405 in MS² spectra, which hinted those maybe the metabolites of ginsenoside Re and ginsenoside Rg₁, by losing of one glucose molecular and/or one rhamnose molecular. By comparison with literature data [35, 37], we suggested that both of them were 20(S) (R)-ginsenoside Rh₁/ginsenoside F₁. M8 showed a molecular ion at m/z 798 [M-H]⁻ in MS spectra, and exhibited m/z 717 [M-H-SO₃]⁻ in MS² spectra, which was consistent with the fragmentation of salvianolic acid B sulfates. In accordance with the literature data on the characteristic of MS/MS [36], M8 was identified as salvianolic acid B sulfates. M9 showed a molecular ion at m/z 783 [M-H]⁻ in MS spectra, and exhibited m/z 621 [M-H-Glu]⁻ and 459 [M-H-2Glu]⁻ in MS² spectra. The results showed the same fragmentation pathway as the metabolite of ginsenoside Rb₁ and ginsenoside Rd. By comparison with literature data [38, 39], M9 was suggested as ginsenoside Rg₃.

Conclusion

By analyzing the constituents in rat serum of FTZ based on UPLC-MS technique and serum pharmacokinetics approach, a method for rapid analysis of the potential effective constituents in a Chinese Medicine formula FTZ have been established. In this study, 27 of the prototype constituents and 9 of the metabolites in rat blood after oral administration of FTZ were identified by the UPLC/Q-TOF system, which enhanced the speed and targeting of bioactive constituents analysis.

These results indicated that most of the alkaloids, ginsenosides, and pentacyclic triterpenes could be observed in rat blood through oral administration of FTZ. Meanwhile the salvianolic acid analogues could be converted into metabolites, such as salvianolic acid B sulfates. Our present work on the comprehensive analysis of the FTZ constituents in rat serum suggest that the serum pharmacochemistry study using UPLC–Q–TOF technique offer a rapid and reliable approach for the identification of potential bioactive compositions for complex herb prescriptions. Systemic pharmacokinetic investigation of the constituents in rat serum after oral administration of FTZ is warranted for better understanding the pharmacokinetic basis of the health benefits of FTZ.

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